### FIGURE 1

# Bunes Saste Fibroblest Growth Factor

I ANT TEA TOC CTC TITT CTC TCC TTT TOT TOG TAG ACG ACT TEA GCC TCT GTC CTT 148 THE TAM AGE TEM TOC CCC ACT THE ACC CCT COT CTT THE GTG ATT TAM AGA 143 TI TCA AMO CCT GCT CTG MCA CAG MCT CTT CCT TOG ATT GCA MCT TCT CTA CTT TEG OUT OUR ANC OUC TTC TCC GIT TIG ANA CGC TAG CGG GGA ANA AAT GGG GGA 218 CAA AGT TBA GTT TAA ACT TTT AAA MGT TGA GTC ACG GCT GGT TGC GCA CGA AAA 124 ICC CCG CAG TOT GGA GAA AGC CTA AAC GTG GTT TGG GTG GTG CGG GGG TTG GGC OG DET DAG TIT TOG GOG ATA AGG GOG GOT GGA DCC CAG GGA ATG CCA AAG CCC TOC COC SEC CTC CSA COC SCS CCC CCC SCC CCT CSC CTC TCC CCC SCC CCC SAC 158 TGA DGC COG SCT CCC CSC COG ACT GAT GTC GCS CGC TTG CGT GTT GTG GCC GAA SIG OCC OCC DAA CTC AGA DOC COG CCC CAG AAA ACC CGA DCB AGT AGG DGG CDG CGC 514 GCA GGA GGG AGG AGA ACT GGG GGC GCG GGA GGC TGG TGG GTG TGG GGG GTG GAG ATG TAG AME ATG TGA COC COC GGC GGG GGG GGC AGA TEA GCG GAC GGC TGC CCG COG TTG CAA COG GAT CCC GGG COC TGC AGC TTG GGA GGC GGC TCT CCC CAG CCG GCG TCC GCG GAM ACA CCC ATC TGT GAA CCC CAM STC CCG GCC CGG CTC OCC SCS CAC CAS GGG CCS GCC GAC AGA AGA GCS GCC SAG CGG CTC GAG GCT GGG 999
GCA GCC GGG AGC ATC ACC ACG CTG CCC GCC TTG CCC GAG GAT GGC GGC AGC GGC ALG ALG GLy Ser Ile The The Lee Pro Ale Lee Pro Gle Asp Gly Gly Ser Gly CCC TTC CCG CCC CGC CAC TTC AMG GAC CCC AMG CGG CTG TAC TGC AMA AMC CGG Ale Phe Fre Fre Gly Ble Phe Lye Asp Pre Lye Arg Lee Tyr Cye Lye Ass Gly 18 GGC TTC TTC CTG CGC ATC CAC CCC GAC GGC GGA GTT GAC GGG GTC CGG GAG AAG Bly Pho Pho Lew Arg Tie Bio Pro Asp Gly Arg Vol Asp Gly Vol Arg Glw Lys 46 AGC GAC CCT CAC ATC AND CTA CAA CTT CAA GCA GAA GAG AGA GGA GTT GTG TCT Set Amp Pro Bis Ile bye Low Gim Low Gim Ale Glu Glu Are Gly Vel Vel Set ATC ANA GEA GTG TGT GCT ANC CGT TAC CTG GCT ATG ANG GAA GAT GGA AGA TTA The Lys Gly Vel Cys Ale Age Arg Tyr Lew Ale ETT Lys Glu Aep Gly Arg Lew 1269 1276 GAT TOT AAA TOT STT ACG GAT GAG TOT TTC TIT TIT GAA CGA TTG GAA TCT Lou Alle Ber Lye Cye Vel The App 118 Cye Phe Phe Sie Acg Lou Giu Ser ANA CGA ACT GGG CAG TAT ANA CTT GGA TCC ANA ACA GGA CCT GGG CAG ANA GCT Lys Arg The Gly Gla Tyr Lys Lee Gly Ser Lys The Gly Fre Gly Gla Lys Ale ATA STE TIT CTT CCA ATG TCT GCT. ANG AGC TGA TIT TAA TGG CCA CAT CTA ATC Ilu Lou Pho Lou Pto AET Set Ale Lys Set 1912
TEA TIT CAC ATG AAA GAA GAA GTA TAT TIT AGA AAT TIG TEA ATG AGA GTA AAA 1519 1566
SAA AAT AAA TGT GTA TAG CTC AGT TTG GAT AAT TGG TCA AAC AAT TTT TTA TCC 1593 LGT AMA ATA TOT AMC CAT GCC CMG TAM MEM AMA ATA ACA AMA GTT GTA MAM TOT ATA TTC TCC CTT TTA TAT TGC ATC TGC TGT TAC CCA GTG AND CTT ACC TMG 1784 1784 CAT CAT THE CAC GCA THE GCT TEA THE GAA AMS AGG CTE TEA AAA TGE LYSS LYST TEA GAA AAC AAA ATT TCT TCA TGG AAA TCA TAT ACA TTA GAA AAT CAC 1839 1836 AGT CAG ATG TIT AAT CAA TCC AAA AAT GTC CAC TAT TTC TTA TGT CAT TCG TTA 1861 127C TAC AND TITL CTA AME ATA TAA AND TOA ANT TAA TCA ANT CCT TYC ANA OTT 1917 144 FFC TCT GGC AGT TCC TTA TGA TAG AGT TTA TAA AAC AGT CCT GTG TAA

1971 ACT GCT GGA AGT TCT TCC GGA ATT C

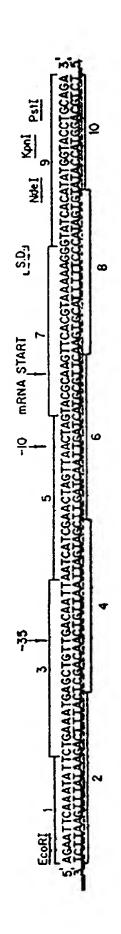
# FIGURE 2 Human Acidic FGF

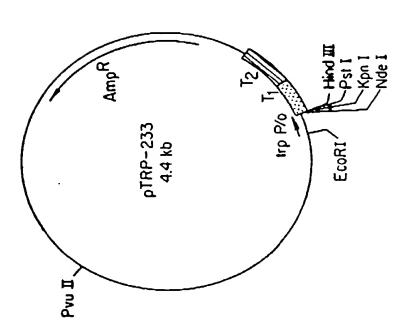
TGC	ATT	TTG	TGC	СТТ	TGC	TGG	AAG	27 AAC	CGA	CTA	CAG	GTT	TGT	TCA	ATT	тст	54 TAC
AGT	СТТ	GAA	AGC	GCC	ACA	AGC	AGC	81 AGC	TGC	TGA	GCC	ATG MET	GCT Ala	GAA Glu	GGG Gly	GAA Glu	108 ATC Ile
ACC Thr	ACC Thr	TTC Phe	ACA Thr 10	GCC Ala	CTG Leu	ACC Thr	GAG Glu	135 AAG Lys	TTT Phe	AAT Asn	CTG Leu	CCT Pro	CCA Pro 20	GGG Gly	AAT Asn	TAC Tyr	162 AAG Lys
AAG Lys	CCC Pro	AAA Lys	CTC Leu	CTC Leu	TAC Tyr 30	TGT Cys	AGC Ser	189 AAC Asn	GGG Gly	GGC Gly	CAC His	TTC Phe	CTG Leu	AGG Arg	ATC Ile 40	CTT Leu	216 CCG Pro
GAT Asp	GGC Gly	ACA Thr	GTG Val	GAT Asp	GGG Gly	ACA Thr	AGG Arg 50	243 GAC Asp	AGG Arg	AGC Ser	GAC Asp	CAG Gln	CAC His	ATT Ile	CAG Gln	CTG Leu	270 CAG Gln 60
CTC Leu	AGT Ser	GCG Ala	GAA Glu	AGC Ser	GTG Val	GGG Gly	GAG Glu	297 GTG Val	TAT Tyr 70	ATA Ile	AAG Lys	AGT Ser	ACC Thr	GAG Glu	ACT Thr	GGC Gly	324 CAG Gln
TAC Tyr	TTG Leu 80	GCC Ala	ATG MET	GAC Asp	ACC Thr	GAC Asp	GGG Gly	351 CTT Leu	TTA Leu	TAC Tyr	GGC Gly 90	TCA Ser	CAG Gln	ACA Thr	CCA Pro	AAT Asn	378 GAG Glu
GAA Glu	TGT Cys	TTG Leu	TTC Phe 100	CTG Leu	GAA Glu	AGG Arg	CTG Leu	405 GAG Glu	GAG Glu	AAC Asn	CAT His	TAC Tyr	AAC Asn 110	ACC Thr	TAT Tyr	ATA Ile	432 TCC Ser
AAG Lys	AAG Lys	CAT His	GCA Ala	GAG Glu	AAG Lys 120	AAT Asn	TGG Trp	459 TTT Phe	GTT Val	GGC Gly	CTC Leu	AAG Lys	AAG Lys	AAT Asn	GGG Gly 130	AGC Ser	486 TGC Cys
AAA Lys	CGC Arg	GGT Gly	CCT Pro	CGG Arg	ACT Thr	CAC His	TAT Tyr 140	513 GGC Gly	CAG Gln	AAA Lys	GCA Ala	ATC Ile	TTG Leu	TTT Phe	CTC Leu	CCC Pro	540 CTG Leu 150
			TCT Ser		TAA	AGA	GAT	567 C <b>TG</b>	TTC	TGG	GTG	TTG	ACC	ACT	CCA	GAG	594 AAG
ттт	CGA	GGG	GTC	CTC	ACC	TGG	TTG	621 ACC	CAA	AAA	TGT	TCC	СТТ	GA			

# Comparison of amino acid sequence of

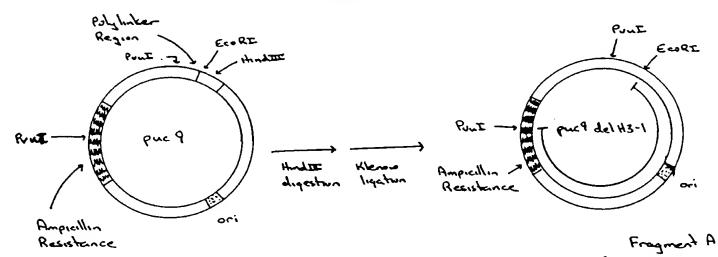
Heparin Binding QLQLSAESVGEVYIKSTETGQYLAMDTDGLLYGSQTPNEECLFLERLEENHYNTYISKKH KLQLQAEERGVVSIKGVCANRYLAMKEDGRLLASKCVTÞECFFFERLESNNYNT|YRSRKY| MAAGSITTLPALPEDGGSGAFPPGH|FKDPKR|LYCKNGGFFLRIHPDGRVDGVREKSDPHI Receptor Binding Domain human basic and acidic FGF AEKNWFVGLKKNGSCKRGPRTHYGQKAILFLPLPVSSD 130—140 TS--WYVALKRTGQYKLGSKTGPGQKAILFLPMSAKS Hepain Cinding (basic/acidic)

Weparin Bindung

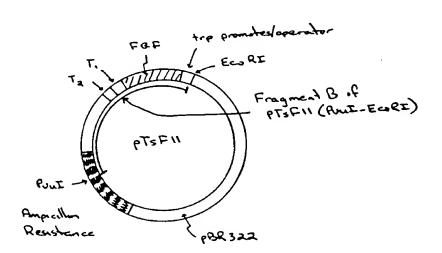


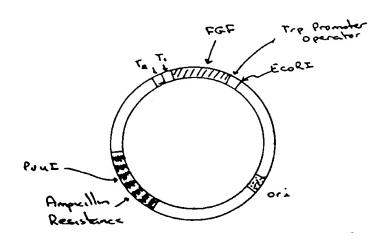


# FIGURE 5



of puc 9 del H3-1 (PULE-ELORE)





Puca del H3 -pTSF-3

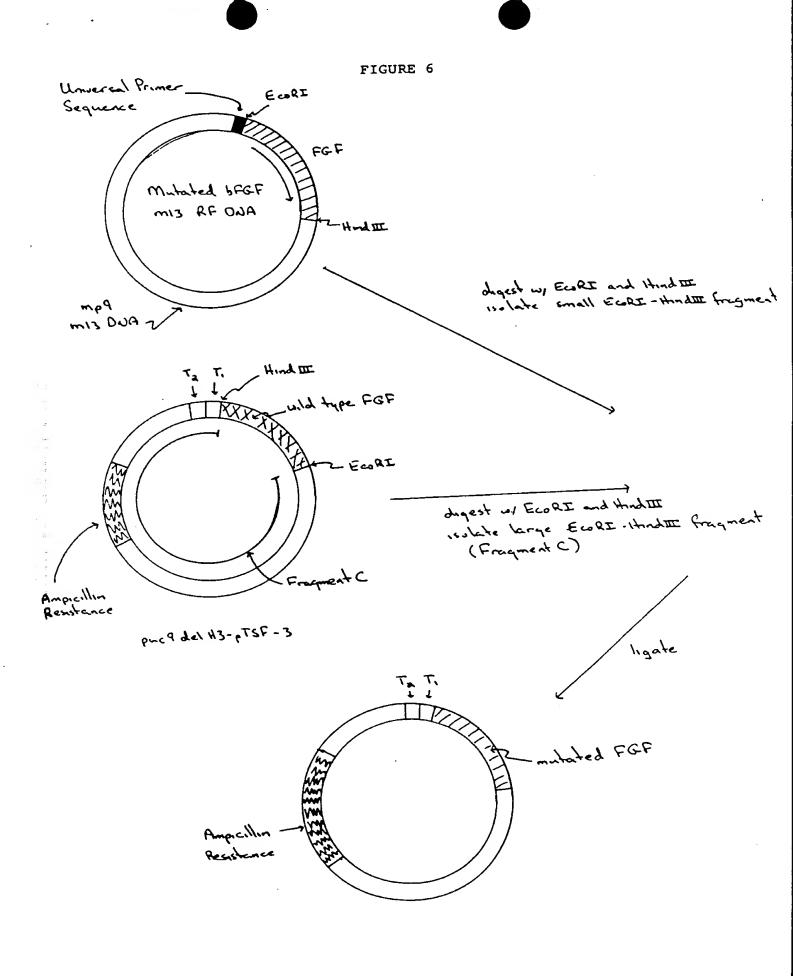


FIGURE 7

